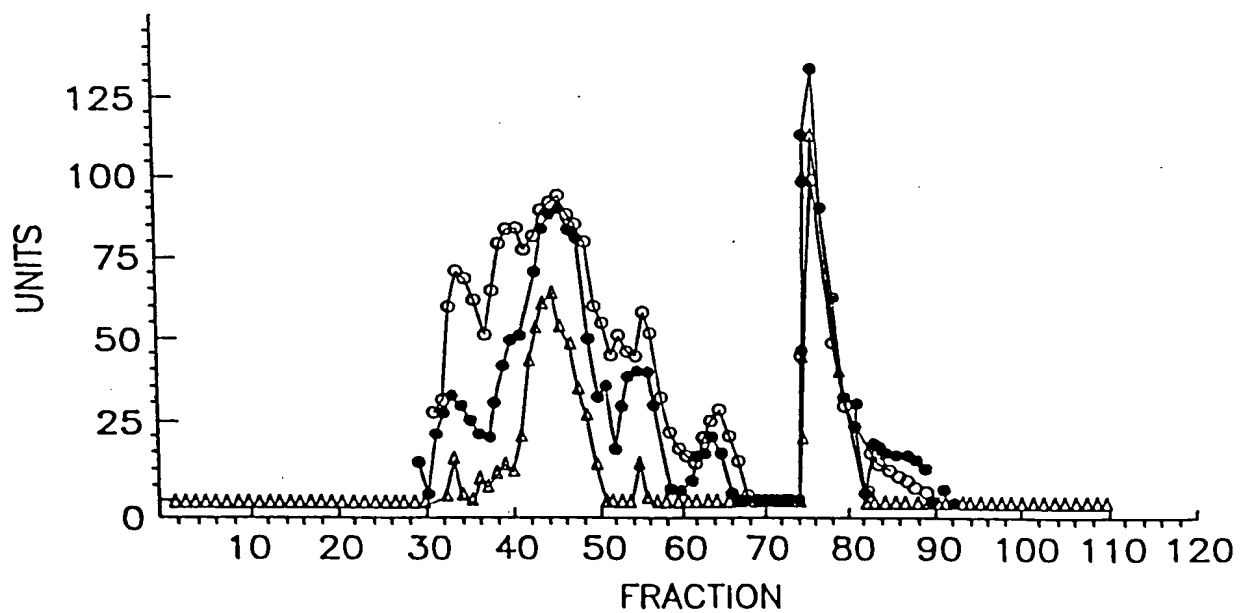


FIG. 1



○ E. COLI
● L. MONOCYTOGENES
△ C. ALBICANS

FIG. 2

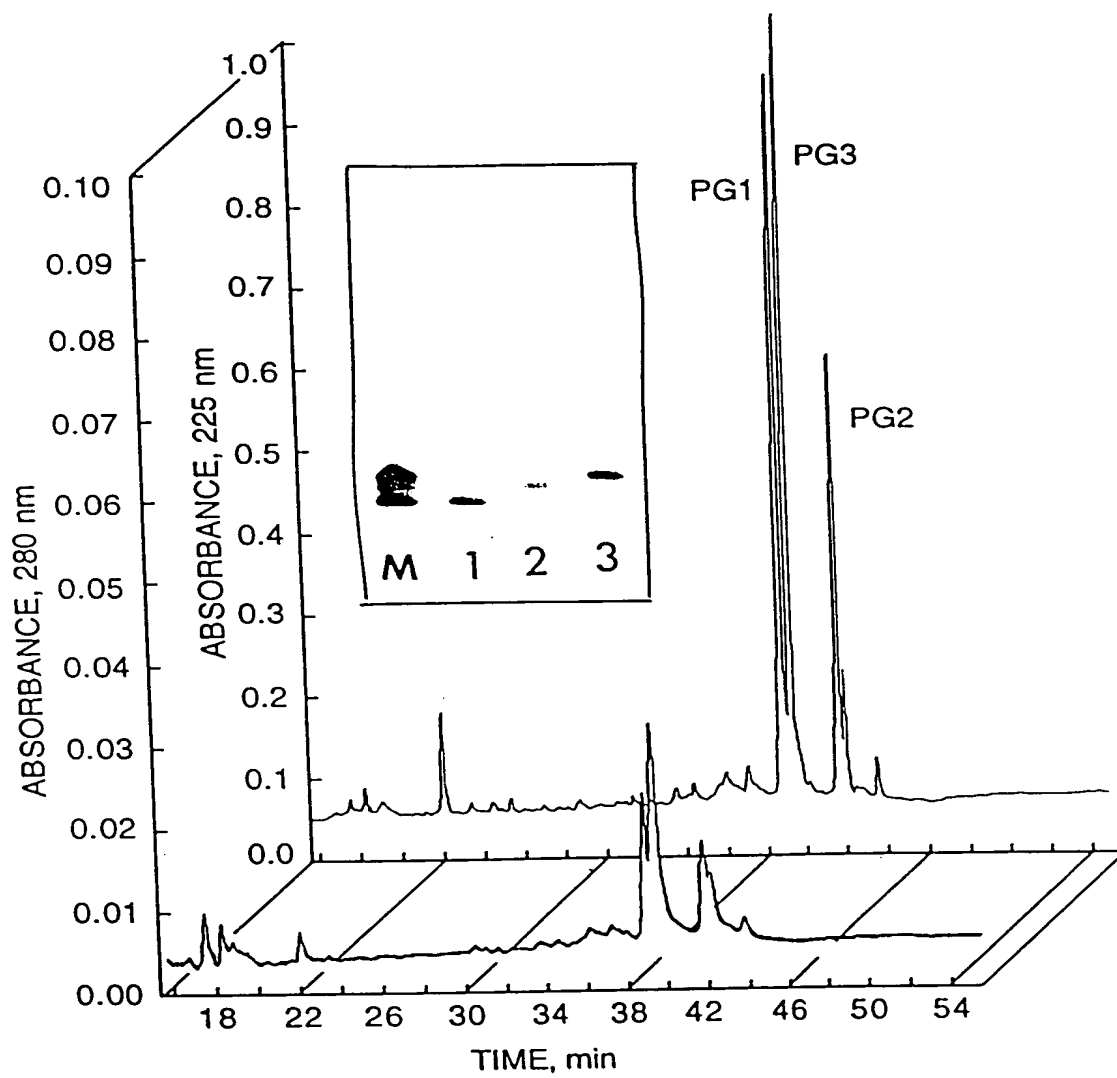


FIG. 3

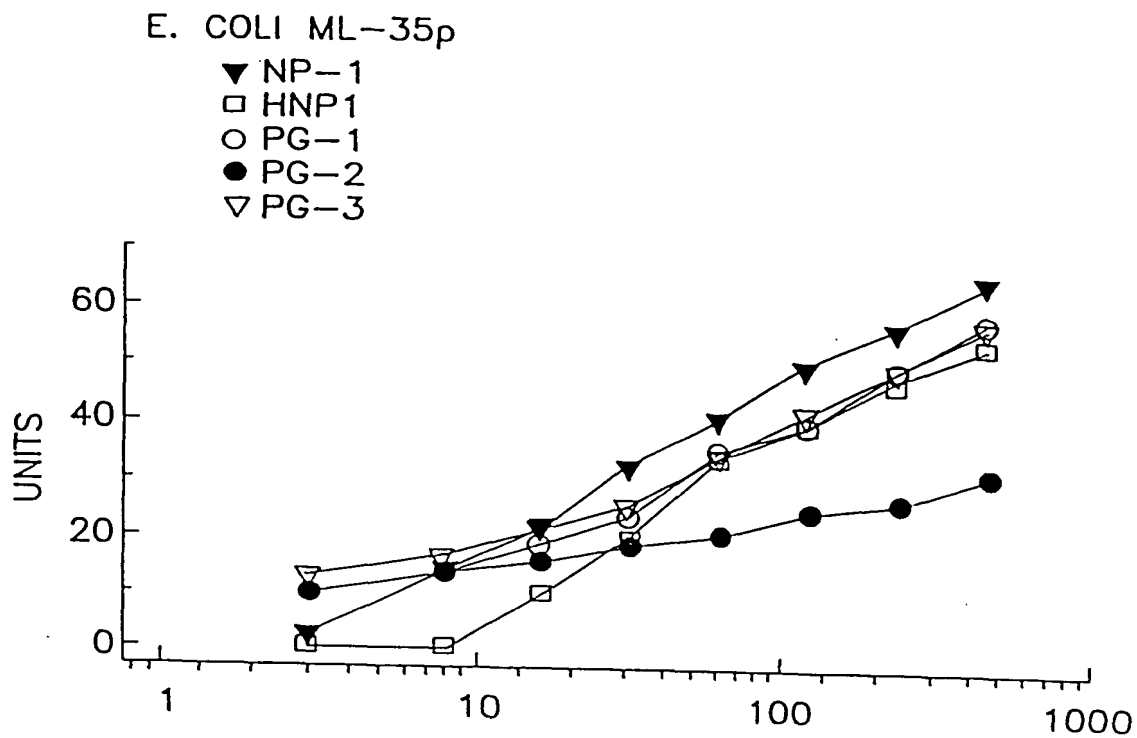


FIG. 4a

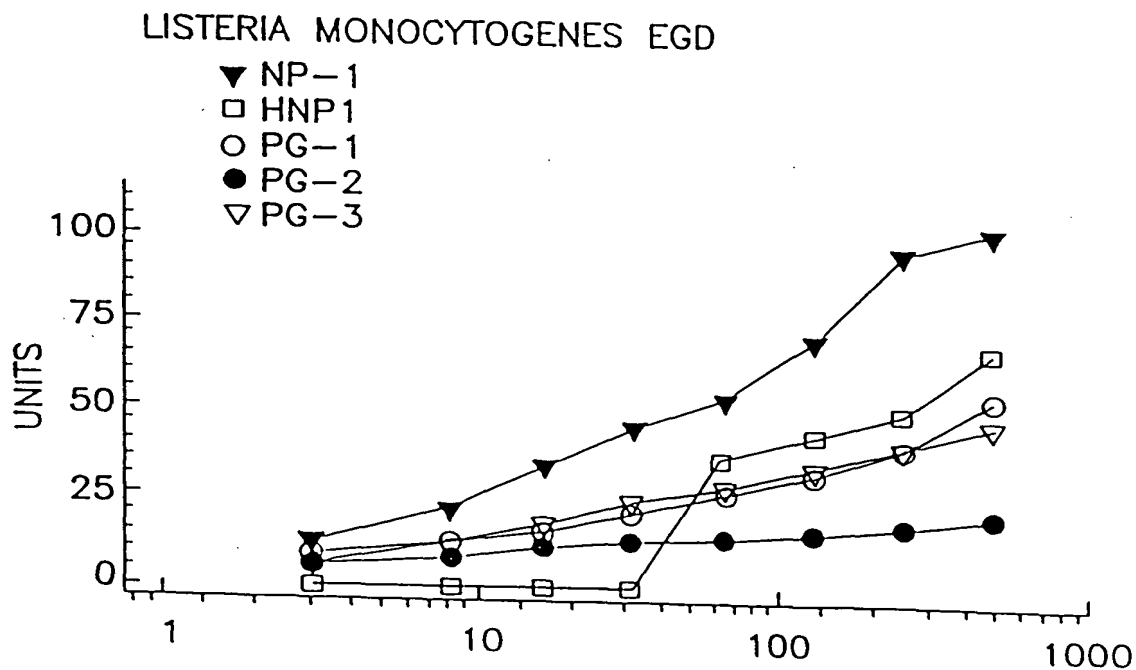


FIG. 4b

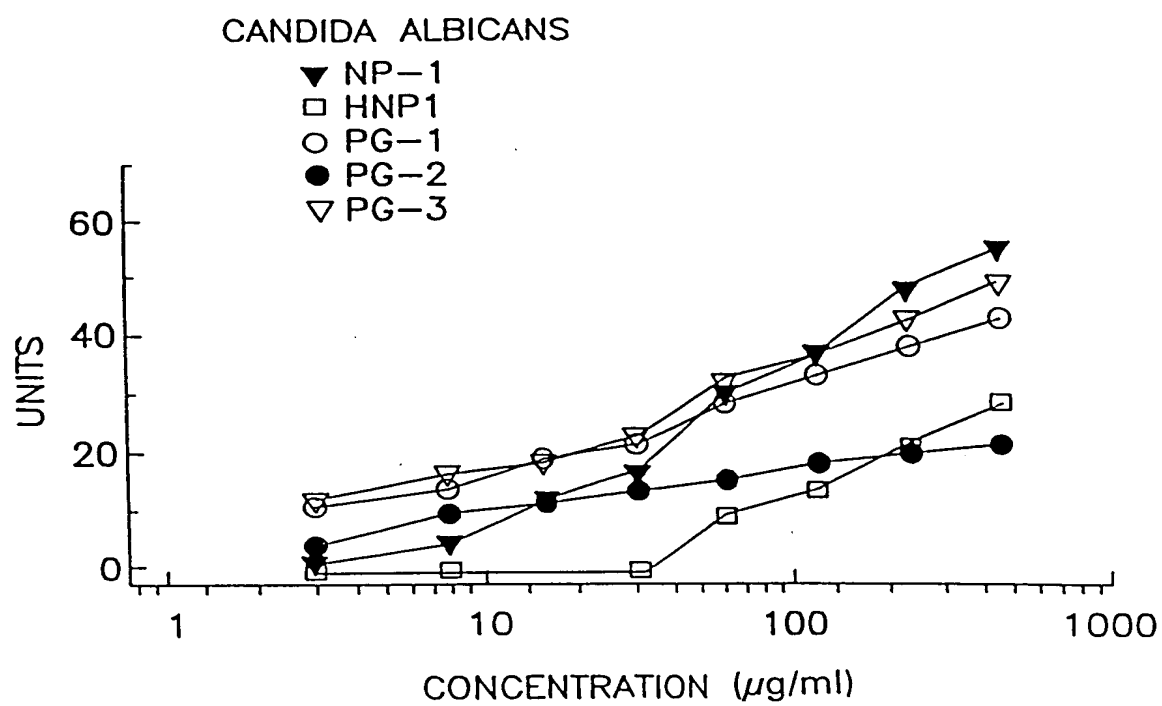


FIG. 4c

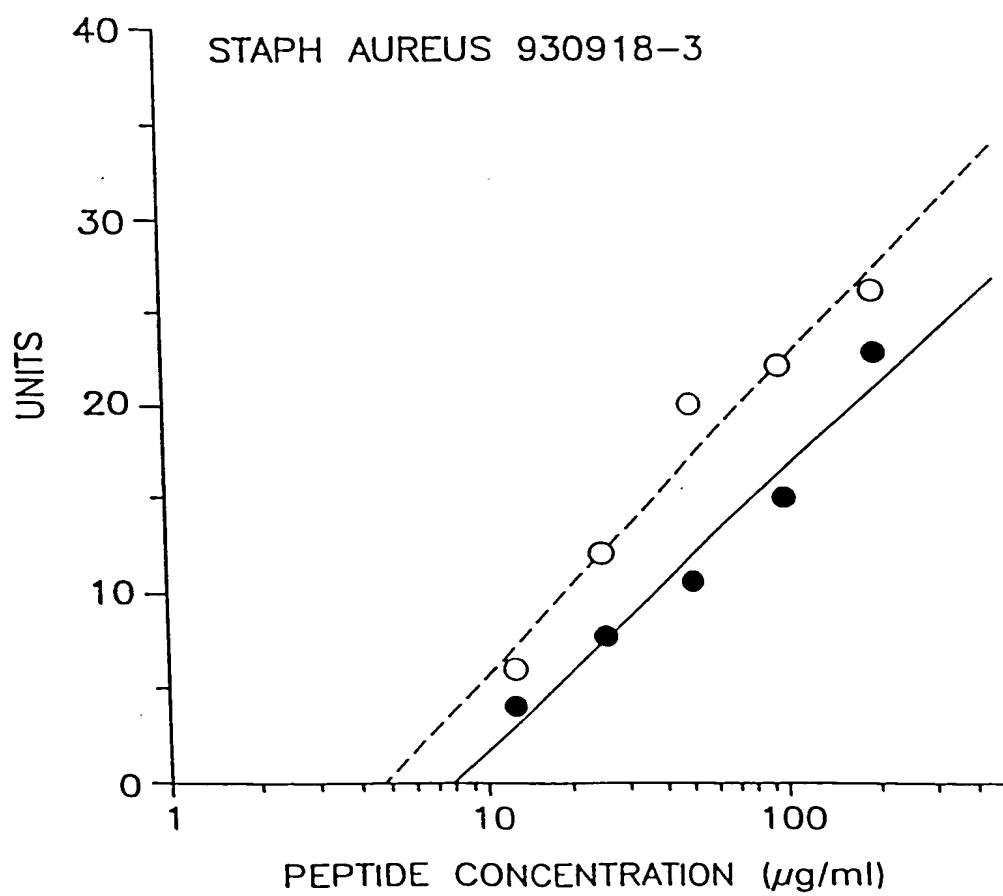


FIG. 4d

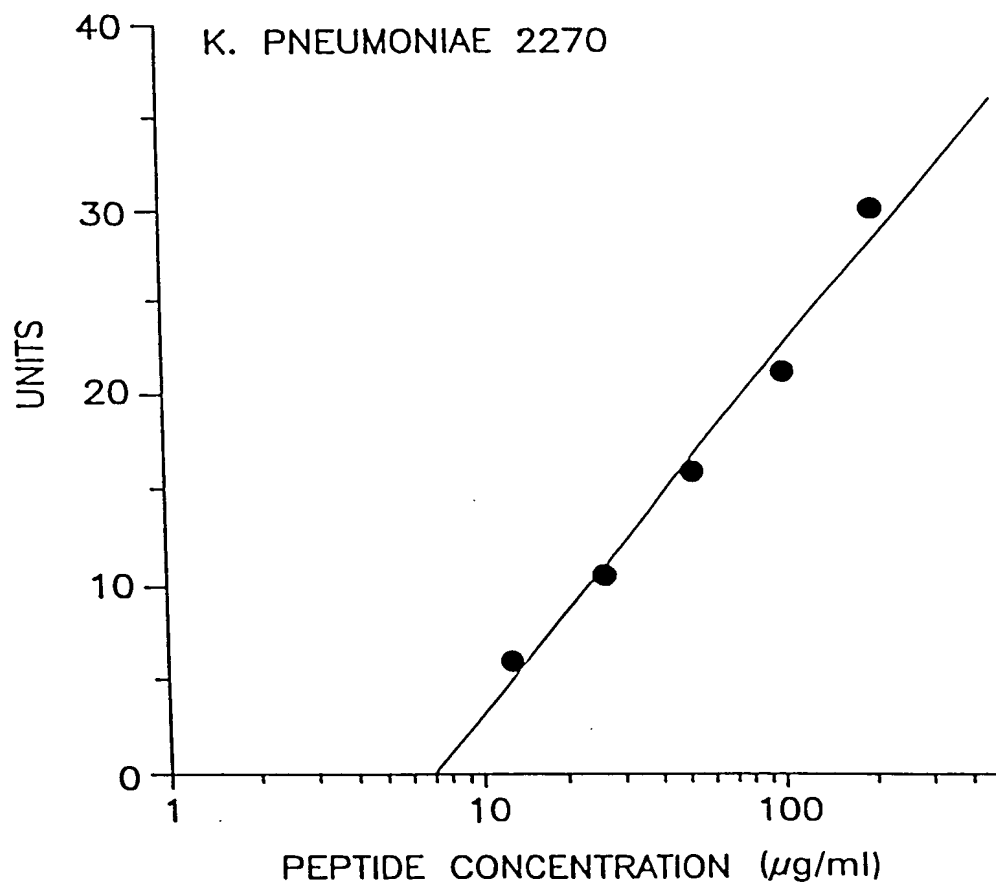
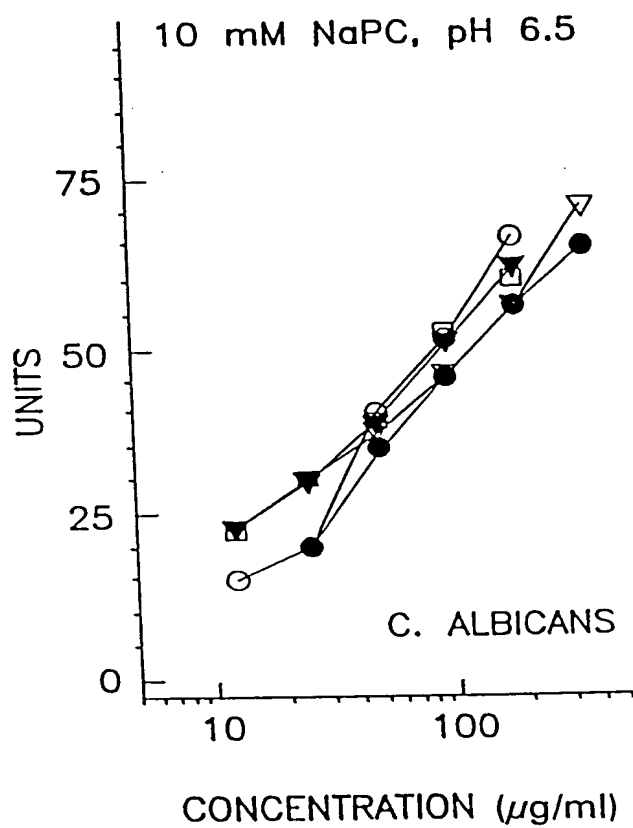
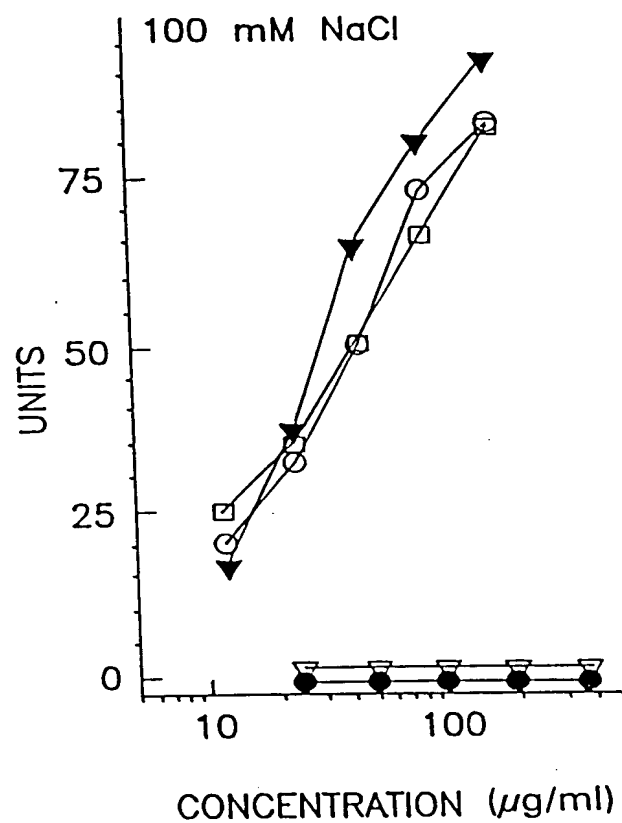


FIG. 4e



○ TP-1
 □ PG-3
 ▼ PG-1
 ▽ NP-1
 ● HNP-1

FIG. 5a-1



○ TP-1
 □ PG-3
 ▼ PG-1
 ▽ NP-1
 ● HNP-1

FIG. 5a-2

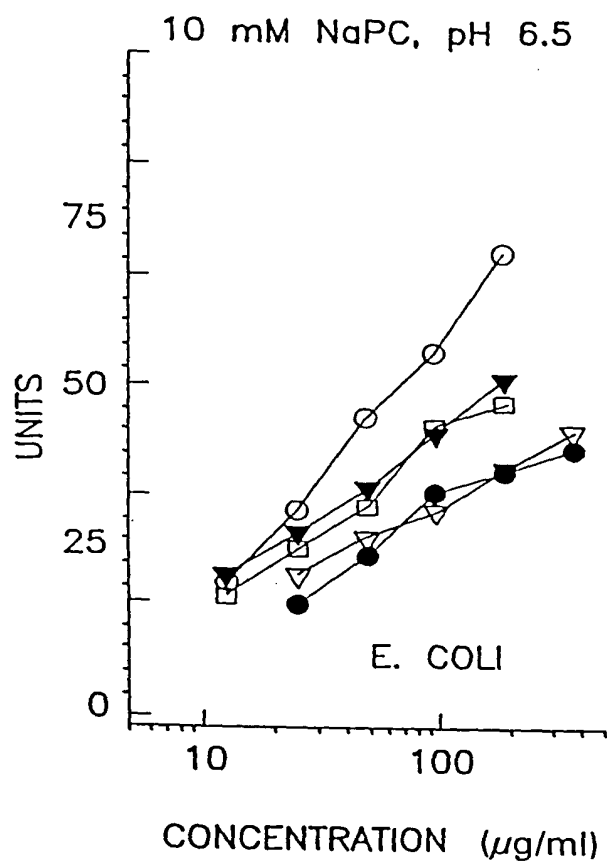


FIG. 5b-1

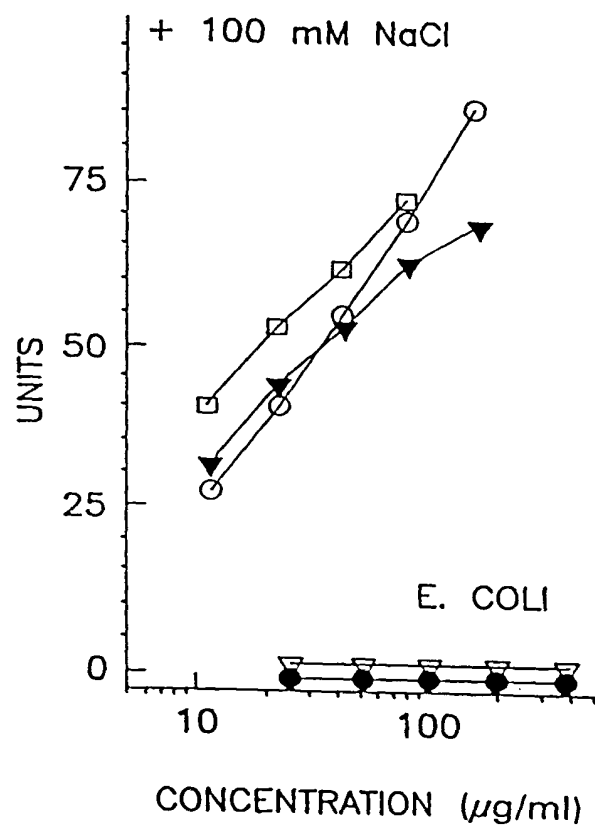
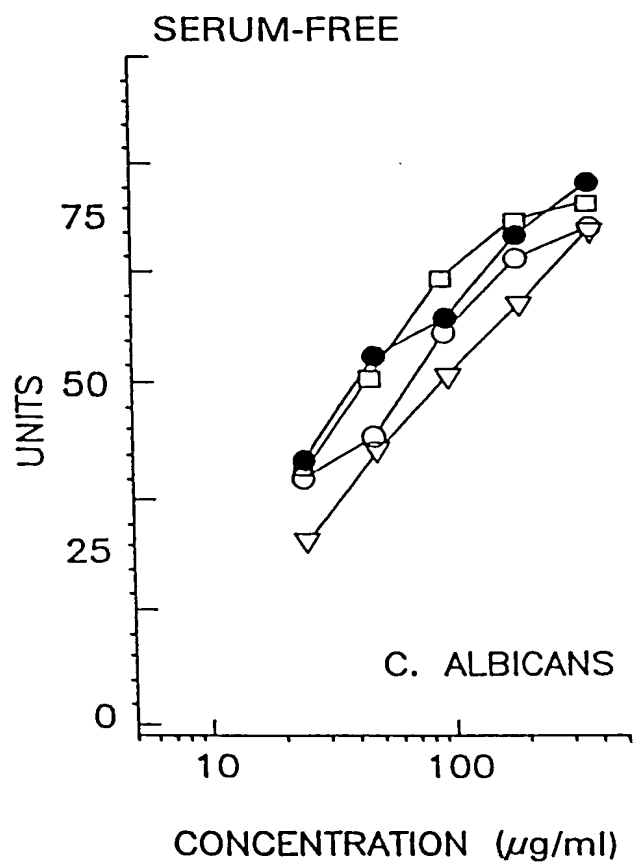
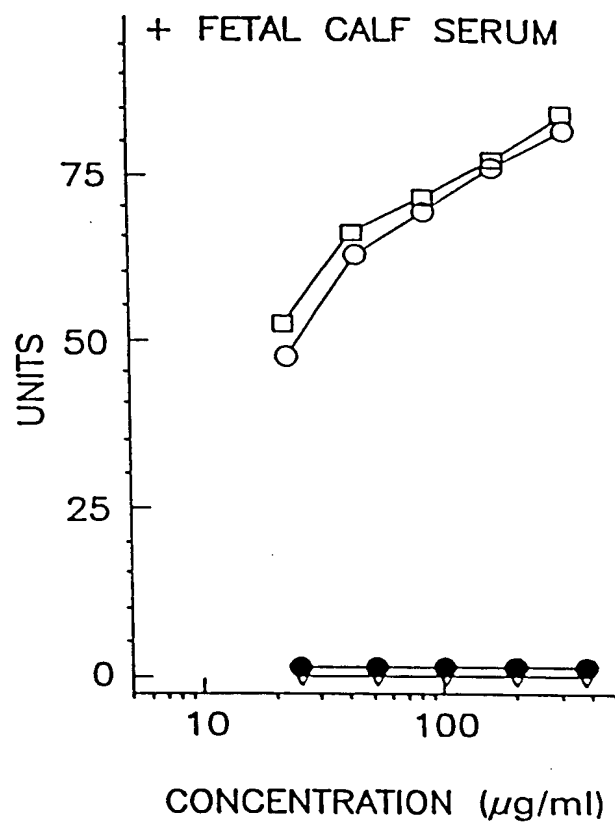


FIG. 5b-2



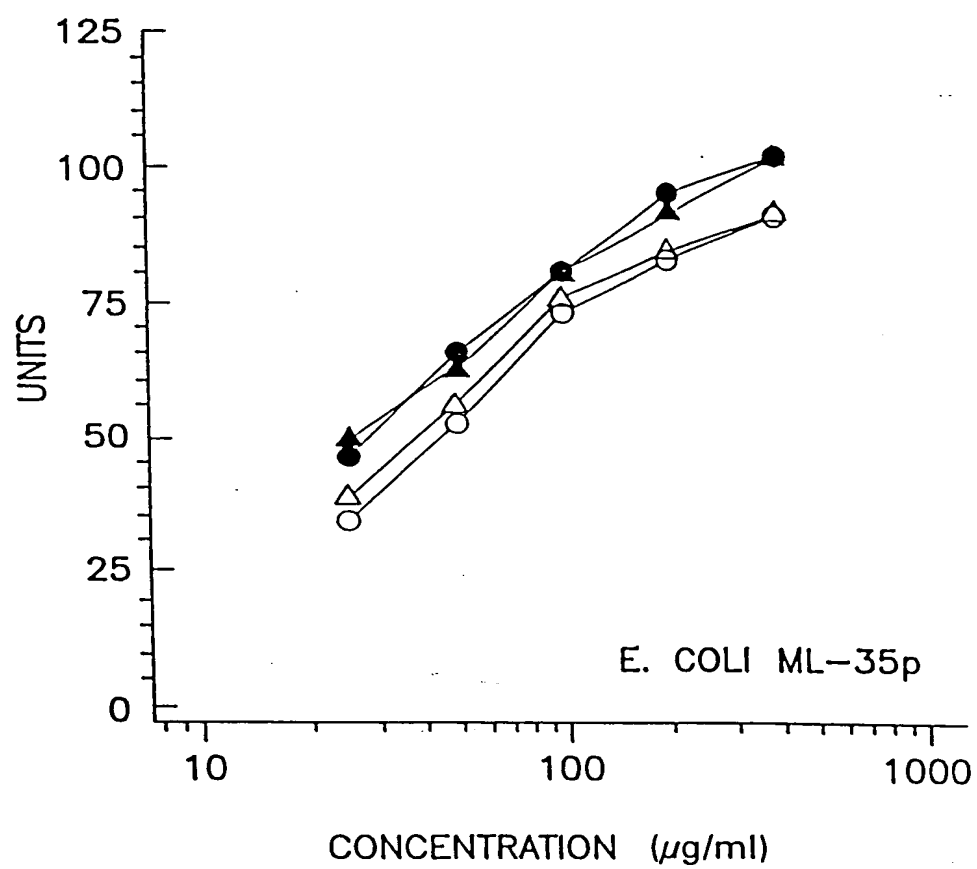
▽ NP-1
○ TP-1
● HNP1
□ PG-3

FIG. 5c-1



▽ NP-1
○ TP-1
● HNP1
□ PG-3

FIG. 5c-2



○ PG-1
● CAM PG-1
△ PG-3
▲ CAM PG-3

FIG. 6a

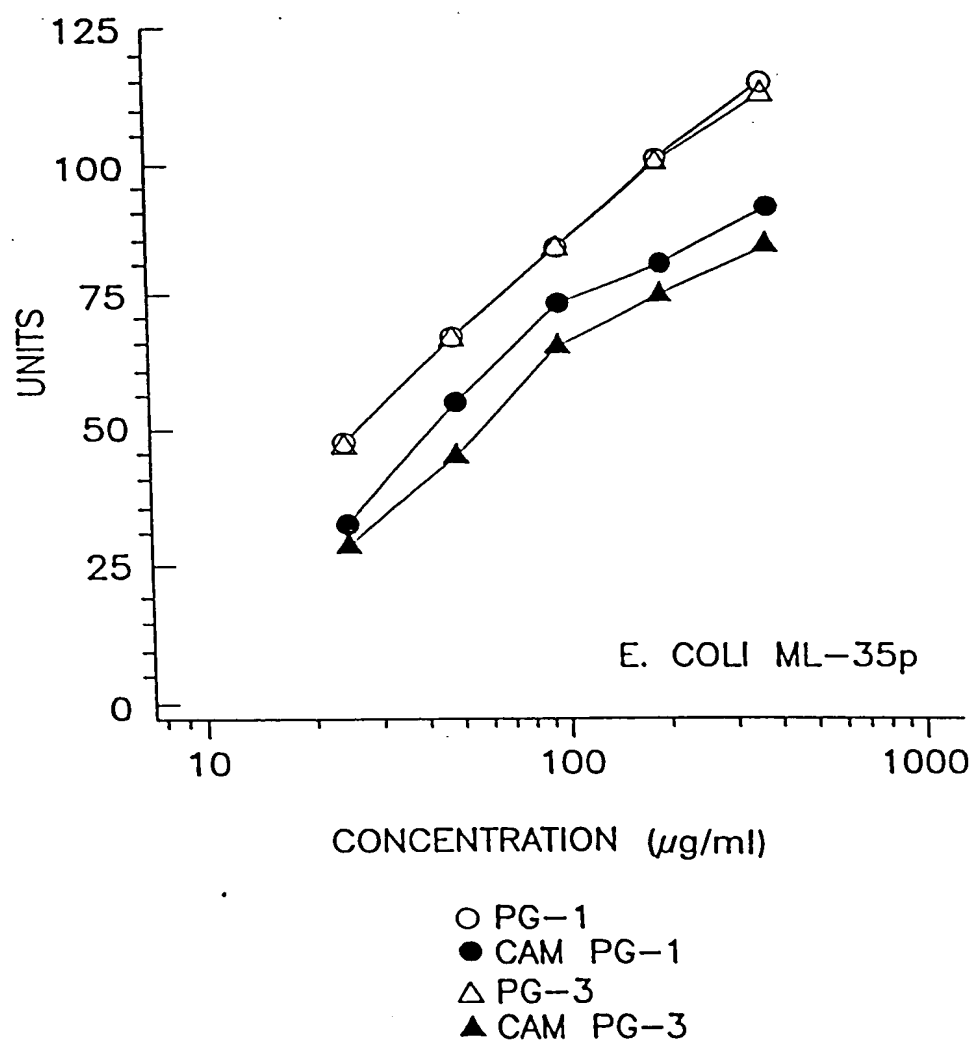


FIG. 6b

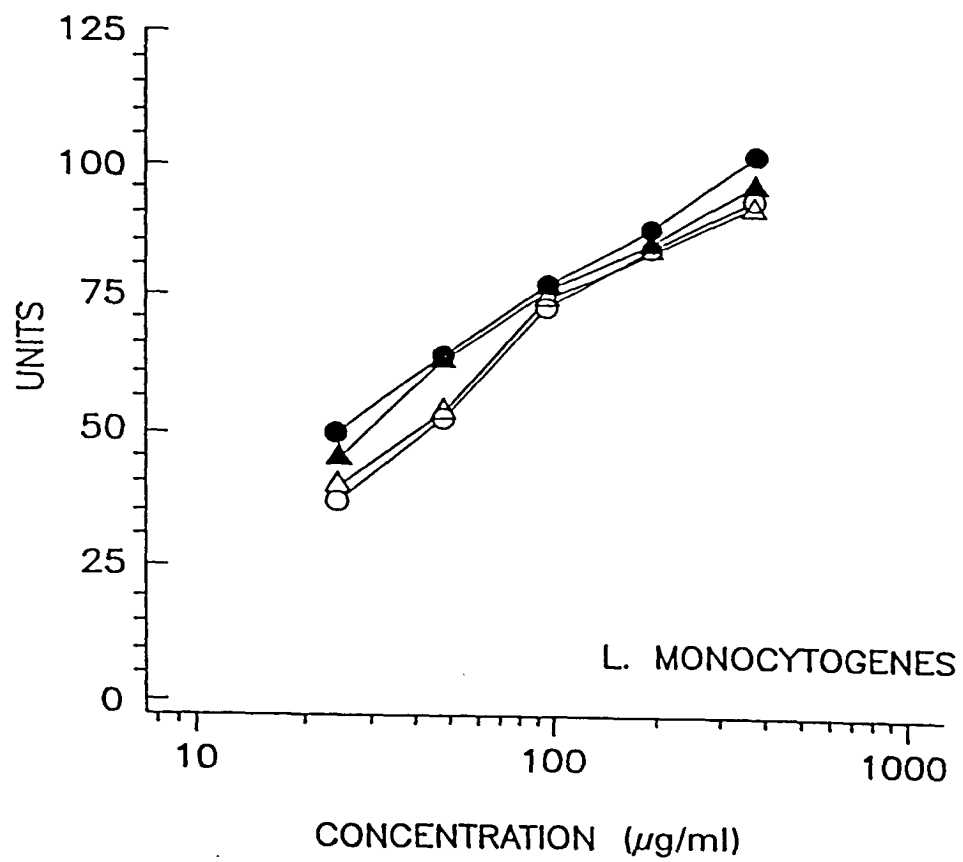


FIG. 6c

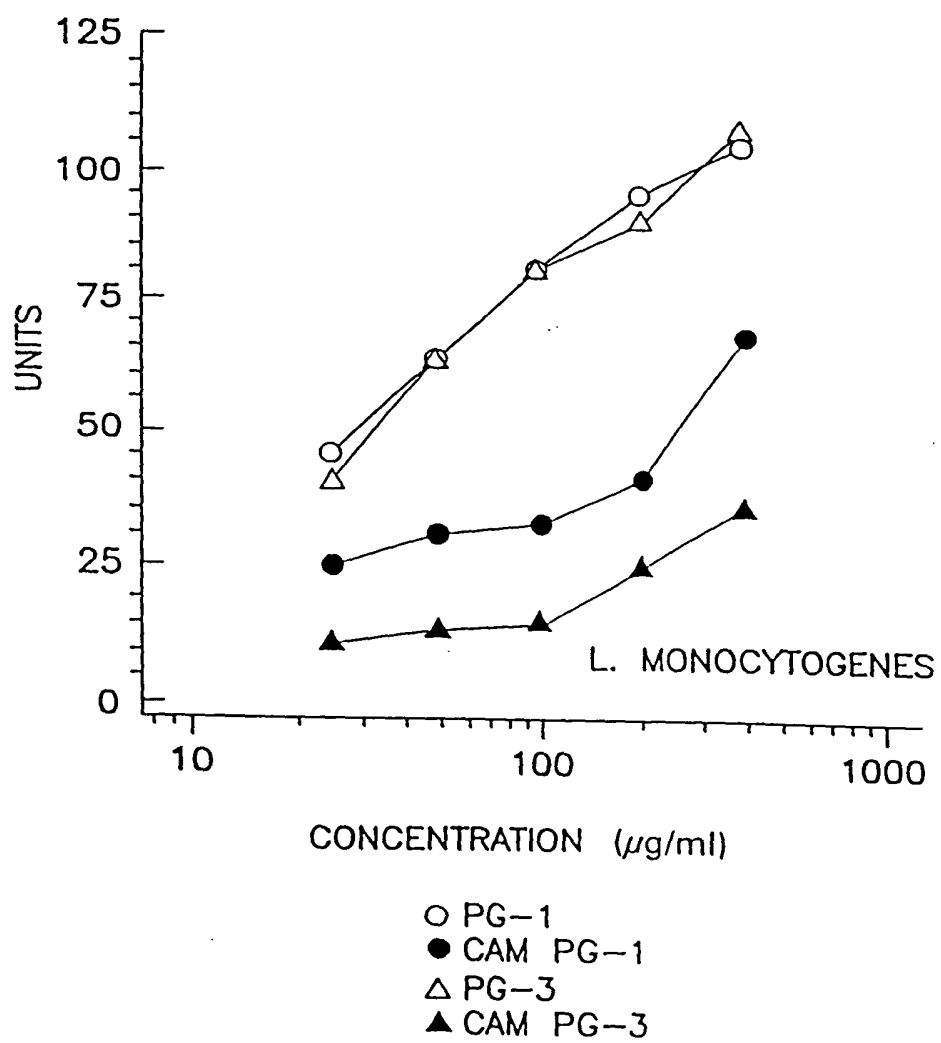


FIG. 6d

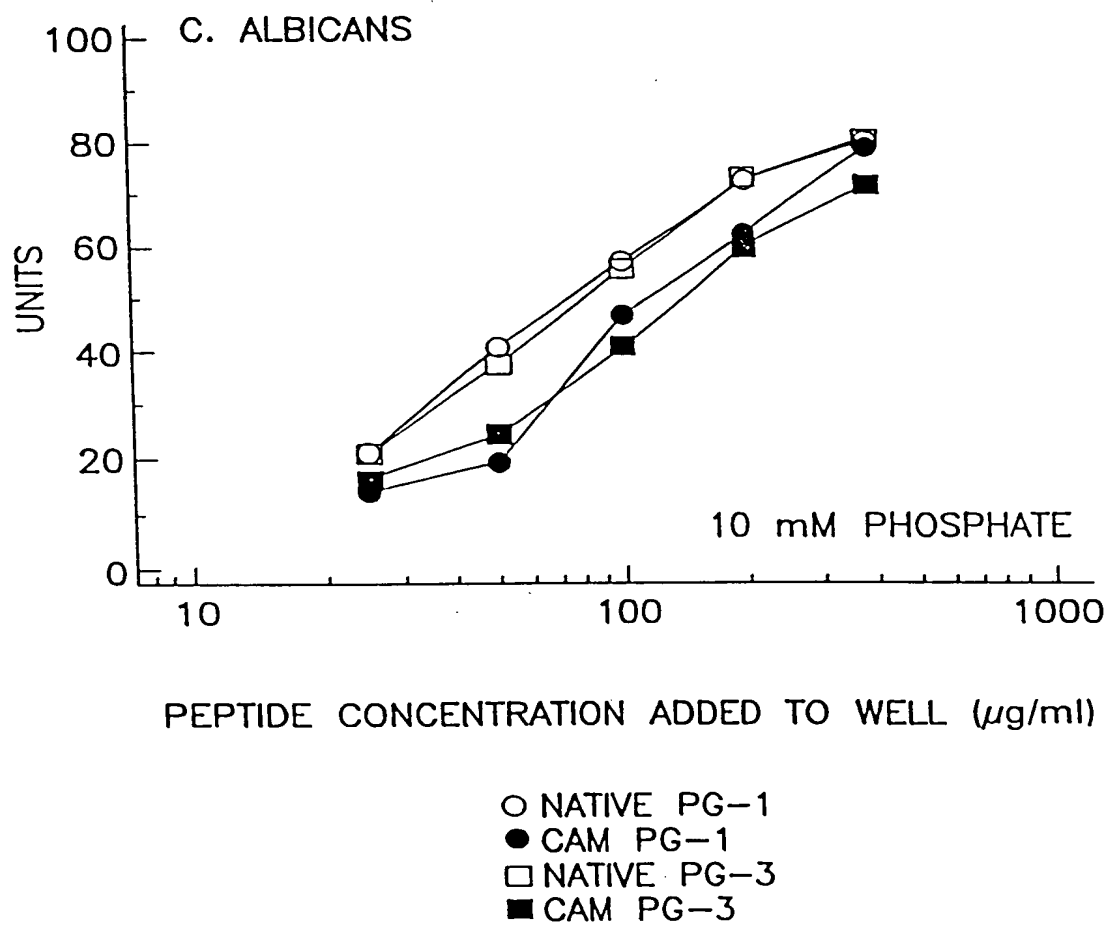
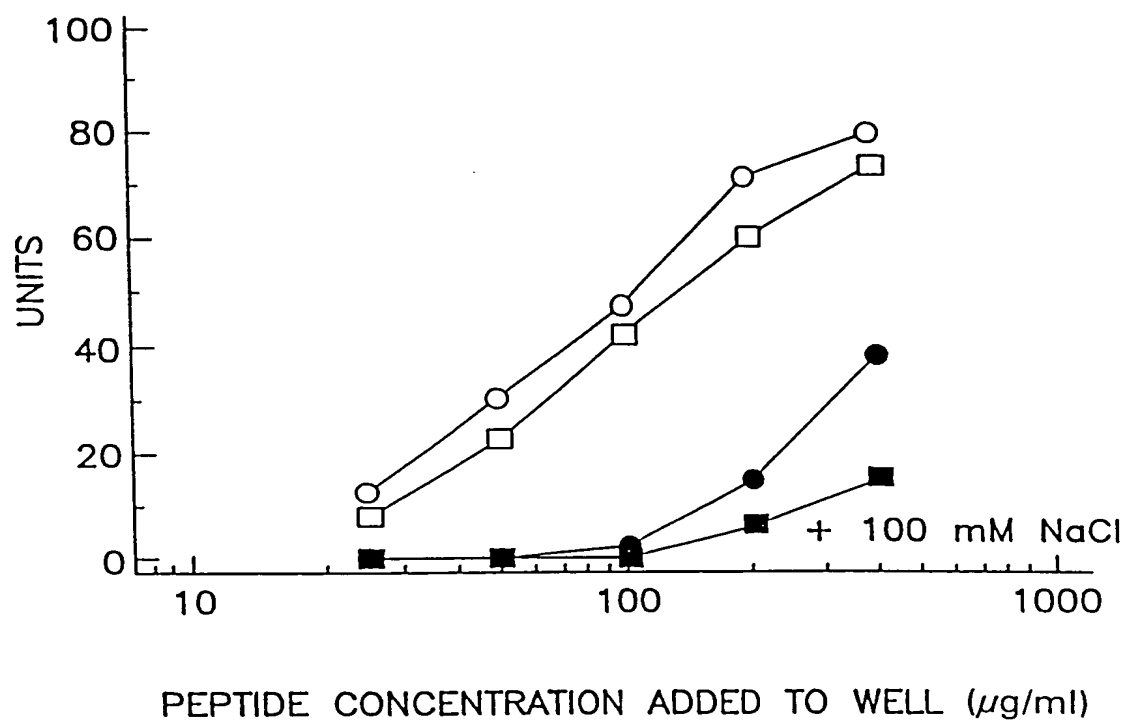


FIG. 6e



○ NATIVE PG-1
 ● CAM PG-1
 □ NATIVE PG-3
 ■ CAM PG-3

FIG. 6f

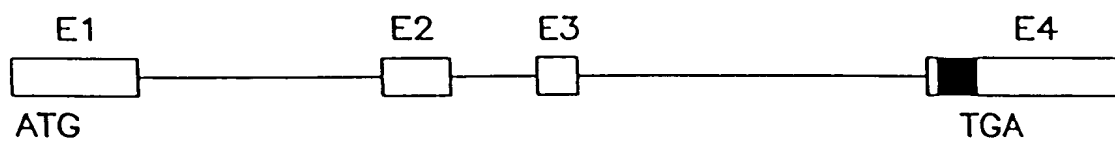


FIG. 9

10	20	30	40	50	
ATGGAGACCGAGAGAGCCAGCCTGTGCCTGGGGCGCTGGTCACTGTGGCTTCTGCTGCTG	60				
MetGluThrGlnArgAlaSerLeuCysLeuGlyArgTrpSerLeuTrpLeuLeuLeuLeu	20				
GCACTCGTGGGTGCCCTCGGCCAGCGCCAGGCCCTCAGCTACAGGGAGGCCGTGCTTCGT	120				
AlaLeuValValProSerAlaSerAlaGlnAlaLeuSerTyrArgGluAlaValLeuArg	40				
GCTGTGGATCGCCTCAACGAGCAGTCCTCGGAAGCTAATCTCTACCGCCTCCTGGAGCTG	180				
AlaValAspArgLeuAsnGluGlnSerSerGluAlaAsnLeuTyrArgLeuLeuGluLeu	60				
GACCAGCCGCCCAAGGCCGACGAGGACCCGGGCACCCCGAAACCTGTGAGCTTCACGGTG	240				
AspGlnProProLysAlaAspGluAspProGlyThrProLysProValSerPheThrVal	80				
AAGGAGACTGTGTGTCCAGGCCGACCCGGCAGCCCCGGAGCTGTGTGACTTCAAGGAG	300				
LysGluThrValCysProArgProThrArgGlnProProGluLeuCysAspPheLysGlu	100				
AACGGGCGGGTGAAACAGTGTGTGGGGACAGTCACCCTGGATCAGATCAAGGACCCGCTC	360				
AsnGlyArgValLysGlnCysValGlyThrValThrLeuAspGlnIleLysAspProLeu	120				
			G ³	G ⁴	
GACATCACCTGCAATGAGGTTCAAGGTGTCAGGGGAGGTGCGCTGTGCTATTGTAGGCGT	420				
AspIleThrCysAsnGluValGlnGlyValArgGlyGlyArgLeuCysTyrCysArgArg	140				
			Gly ³	Gly ⁴	
T ⁴ A ⁴ A ²					
T ⁴ T ⁴ T ²					
AGGTTCTGCGTCTGTGTGTCGGACGAGGATGACGGTTGCGACGGCAGGCTTTCCTCCCCCA	480				
ArgPheCysValCysValGlyArgGly---	149				
Trp ⁴ Ile ⁴ Phe ⁴					
Ile ² ---2					
ATTTTCCCGGGGCCAGGTTTCCGTCCCCCAATTTTCCGCCTCCACCTTTCGGCCCCGCA	540				
			A ² G ²		
CCATTCGGTCCACCAAGGTTCCCTGGTAGACGGTGAAGGATTTCAGGCAACTCACCCAG	600				
			C ⁴		
AAGGCCTTTCGGTACATTAAATCCCAGCAAGGAGACCTAAGCATCTGCTTTGCCCAGGC	660				
CCGCATCTGTCAAATAAATTCTTGTAACC	691				

FIG. 7

ATGGAGACCCAGAGAGCCAGCCTGTGCCTGGGGCGCTGGTCACTGTGGCTTCTGCTGCTG	60
<u>M E T Q R A S L C L G R W S L W L L L L</u>	
^{G5} GCACTCGTGGTGCCCTCGGCCAGCGCCCAGGCCCTCAGCTACAGGGAGGCCGTGCTTCGT	120
<u>A L V V P S A S A Q A L S Y R E A V L R</u>	
^{G5} GCTGTGGATCGCCTCAACGAGCAGTCCTCGGAAGCTAATCTCTACCGCCTCCTGGAGCTG	180
<u>A V D R L N E Q S S E A N L Y R L L E L</u>	
GACCAGCCGCCCAAGGCCgtgagtcgggcaggggctcaggaggggctggggggcgggggc	240
<u>D Q P P K A</u>	
tgtccccacccgccccggggctccctgtccctccccctgctcaggctgtccctcctgcc	300
aggaaggcacttgtccctctaagggggacccctctgccaaggaaaccttcccagagctgg	360
gtgccc tggccgcgtgagagcttcccgcc ttagcctctgggctgtgggctcagggccctg	420
cacagcc tgtgaggcaggagcgggc tctgtcccc tcccc tgtgcacccagcaccaagccc . .	480
agggccaggctcccagcaggggctgcagaggctgctgtctagggtgggggcggggaggggg	540
tgacagatccgagggggaagcctgagcccgagtcccatctccccactttgatccttgacc	600
^{A5} agGACGAGGACCCGGGCACCCGAAACCTGTGAGCTTCACGGTGAAGGAGACTGTGTGTC	660
<u>D E D P G T P K P V S F T V K E T V C</u>	
CCAGGCCGACCCGGCAGCCCCGGAGCTGTGTGACTTCAAGGAGAACGGGgtgaggctgg	720
<u>P R P T R Q P P E L C D F K E N G</u>	
gggctggggggcgc tggcggatgcttccc aaggagctgaacaggagagcctgctggggaag	780
atgtccaggccctgggggtgaggctgggagctcatggatggaggaggggggggtcccagttt	840
^{t3} gaccttgagctctcccc ttccagCGGGTGAAACAGTGTGTGGGGACAGTCACCCTGGATCA	900
<u>R V K Q C V G T V T L D Q</u>	
GATCAAGGACCCGCTCGACATCACCTGCAATGAGgtgagtgggcccttattgggtgtcaag	960
<u>I K D P L D I T C N E</u>	
ttgctaattgggttgggtgtggggaacttccttgggagtggttaccgcgtgccccatccagggc	1020
gtggaaaggccctcctaccccggcccttccctcacctcggccccagggtccaggctctgg	1080
ctctgtcatccttagggccgcggttccctcaatgggggtccccccctcgtatttgtcagaa	1140
^{g3,5} aggcacatttcaggccccaccccgacctctgaatcacactcttgggtggagcccagcct	1200
tgtctcttctcccaagatcccagcgggttcttcc tgtgctgtcggctgagaggcagtgac	1260
cggactaatggacttgcaggccc tgc tcttgccagctttgcggggctggggttgggacc	1320
ctggcaaggccccagccatctctgggcc tgagtccacttatgtgtctgtgggggattcaa	1381
^{g3,5} ^{t5} ccacgtgcttccaaagggtcacagccagagggtggaccagggccccaagcctcttactgtttc	1440

FIG. 8a

cccattcaggggatttttctagtcctggagggaaggggttcttgaccttgccagacc 1500
ccacccgaaacctgtttctcttggtcacagGTTCAAGGTGTCAGGGGAGGTCGCCTGTGC 1560
 F Q G V R G G R L C
 G3

TATTGTAGGCGTAGGTTCTGCGTCTGTGTCCGACGAGGATGACGGTTGCGACGGCAGGCT 1620
Y C R R R F C V C V G R G ***
P5
TTCCCTCCCCCAATTTTCCGGGGGCCAGGTTTCCGTCCCCCAATTTTCCGCCTCCACCT 1680
TTCCGGCCCGCACCATTCGGTCCACCAAGGTTCCCTGGTAGACGGTGAAGGATTTGCAGG 1740
 C3,5
CAACTCACCCAGAAGGCCTTTCGGTACATTAAAATCCCAGCAAGGAGACCTAAGCATCTG 1800
CTTTGCCCAGGCCCGCATCTGTCAAATAAATTCTTGTAACC 1843

FIG. 8b

	1 2 3	4	5 6 7 8 9	10 11 12	13	14	15 16	17 18
PG-1	RGG	R	LCYCR	RRF	C	V	CV	GR*
PG-2	RGG	R	LCYCR	RRF	C	I	CV	GR*
PG-3	RGG	G	LCYCR	RRF	C	V	CV	GR*
PG-4	RGG	R	LCYCR	GW	C	F	CV	GR*
PG-5	RGG	R	LCYCR	PRF	C	V	CV	GR*

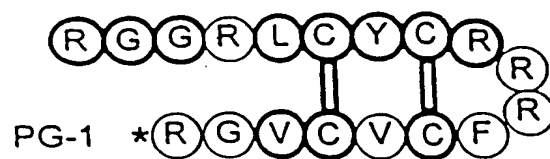


FIG. 10

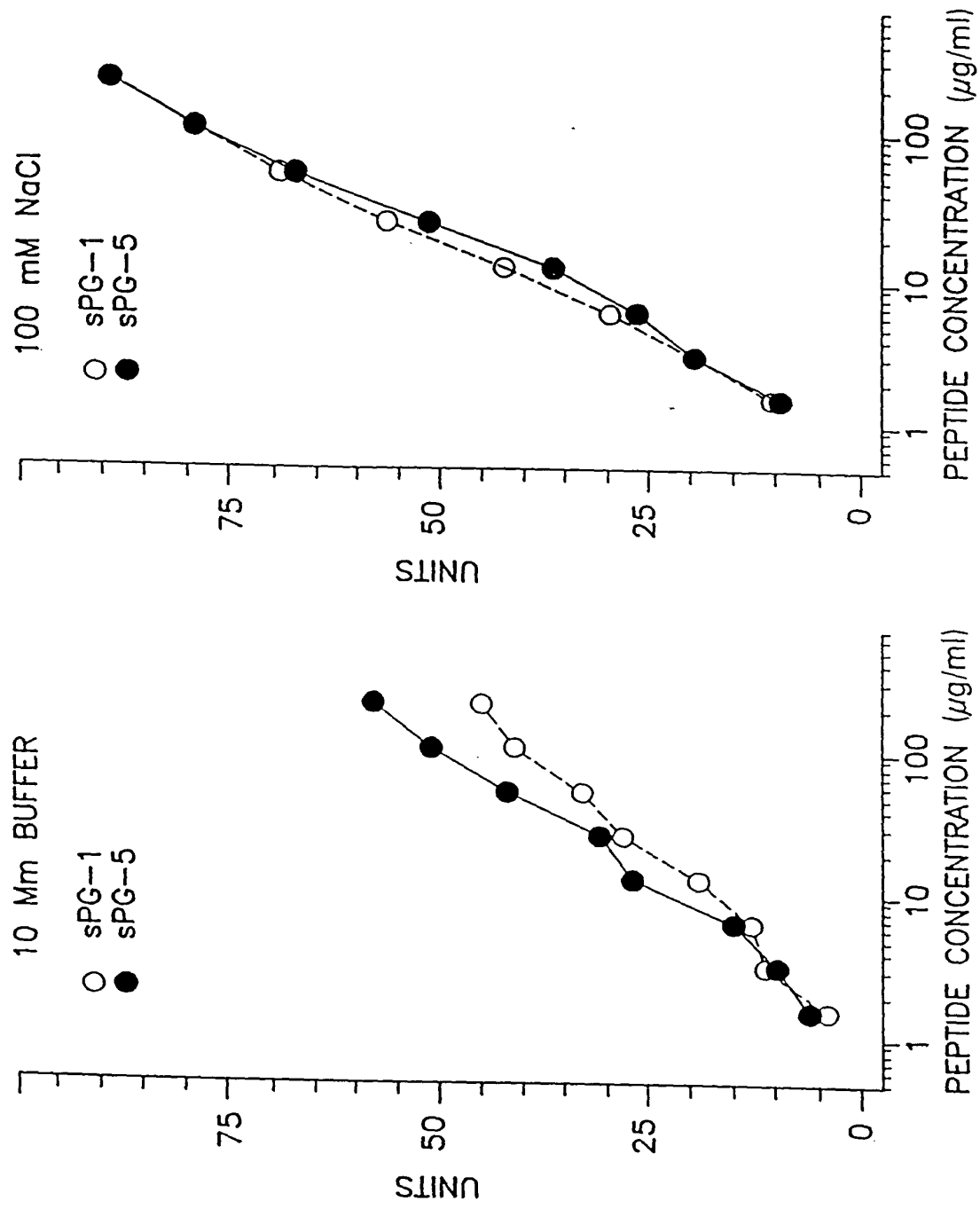


FIG. 11a-1

FIG. 11a-2

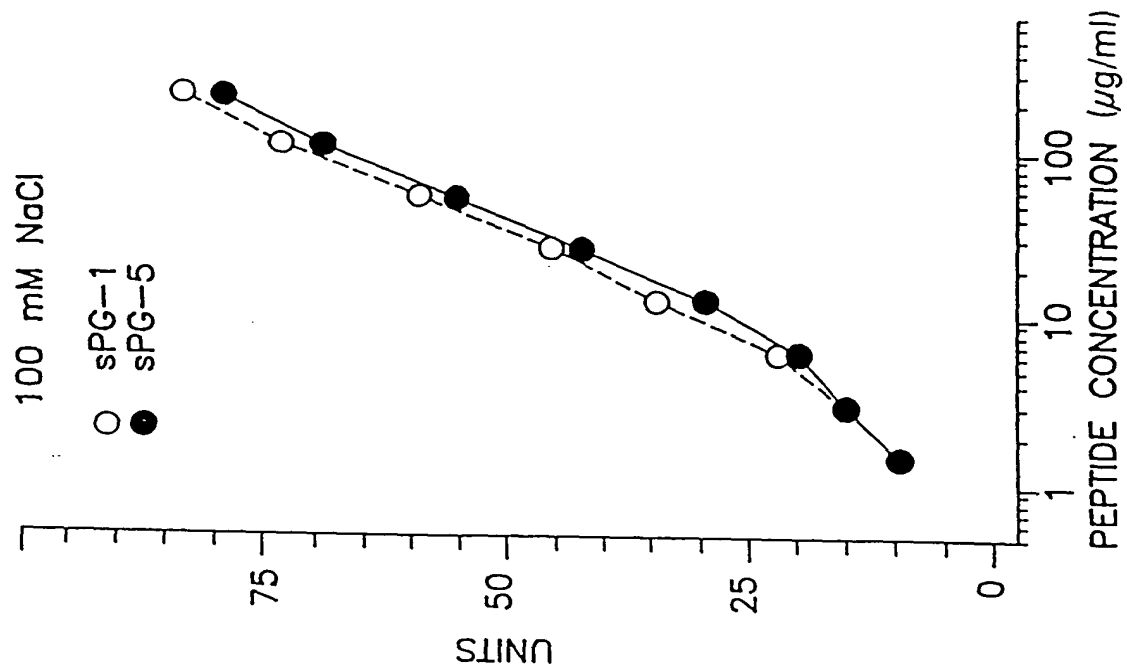


FIG. 11b-2

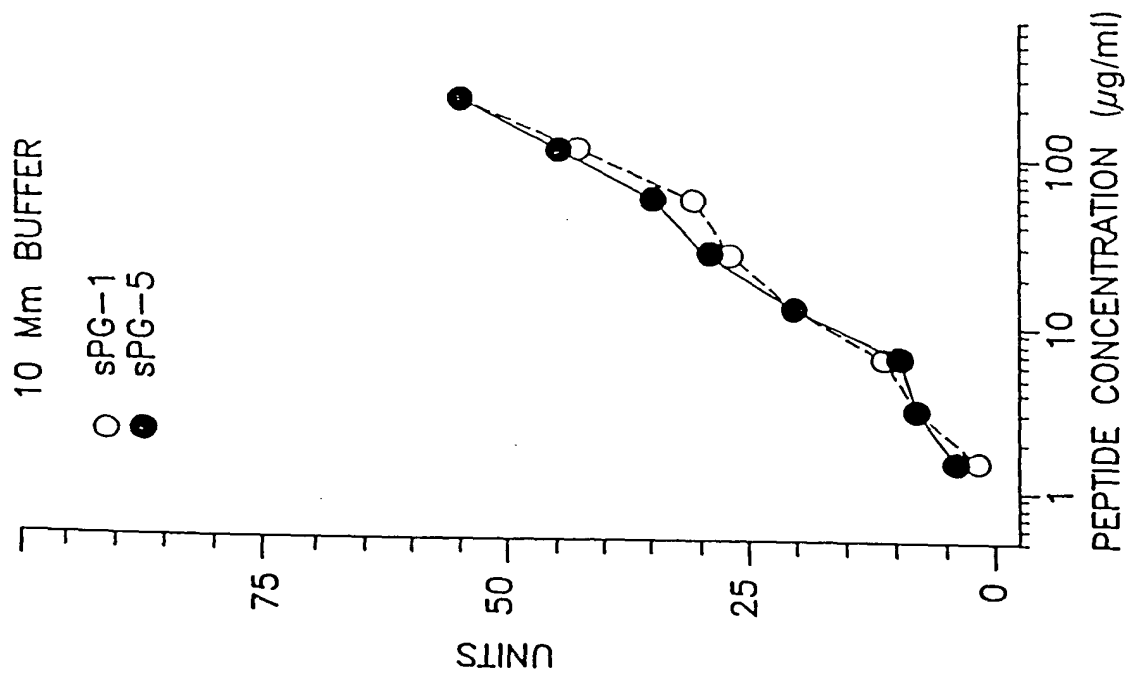


FIG. 11b-1

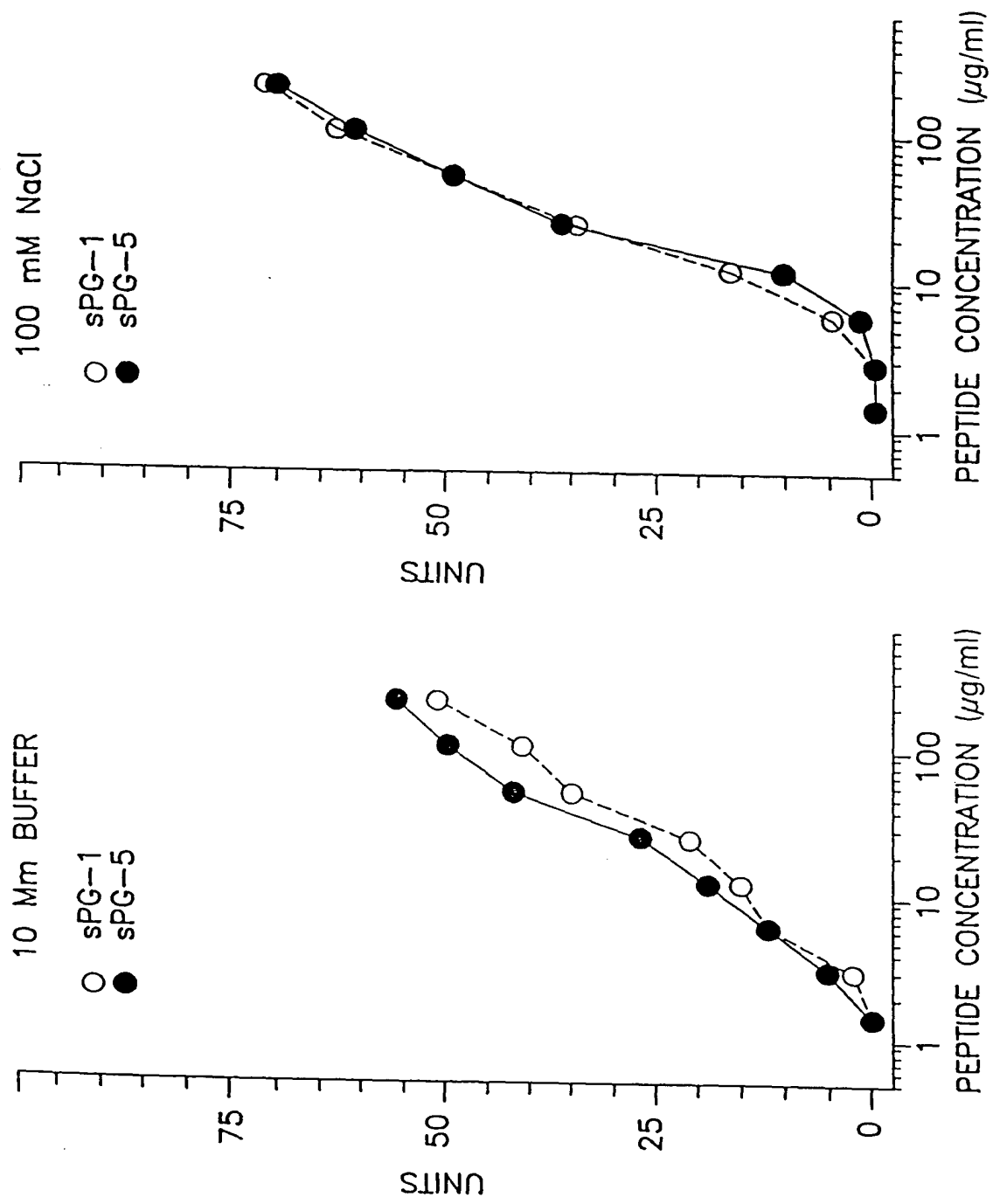


FIG. 11c-1

FIG. 11c-2

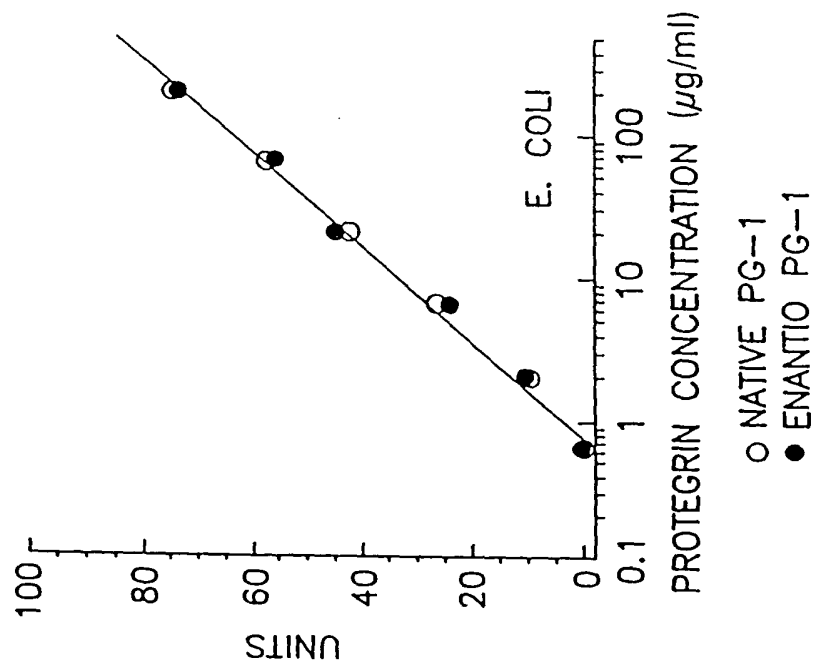


FIG. 12a

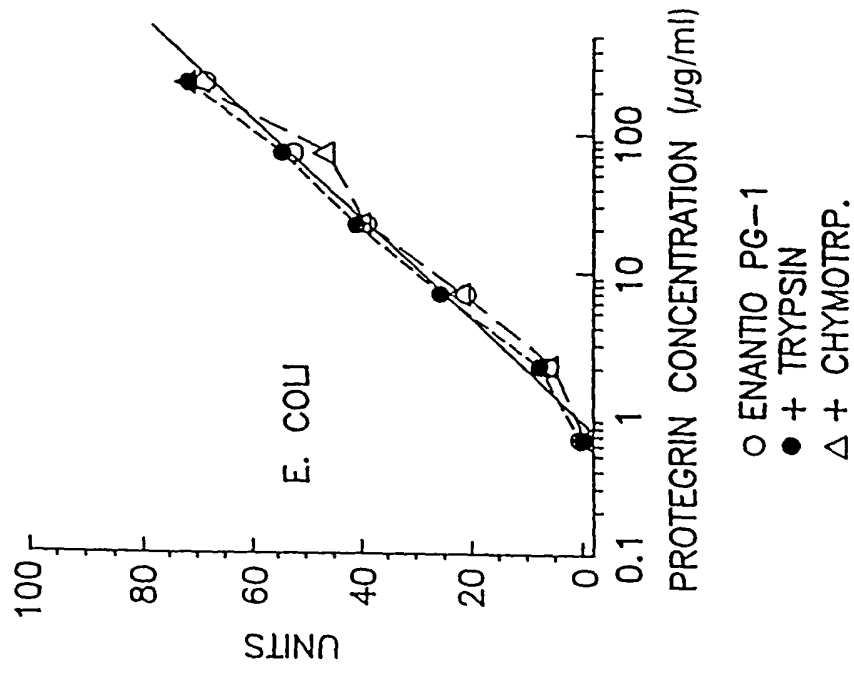


FIG. 12b

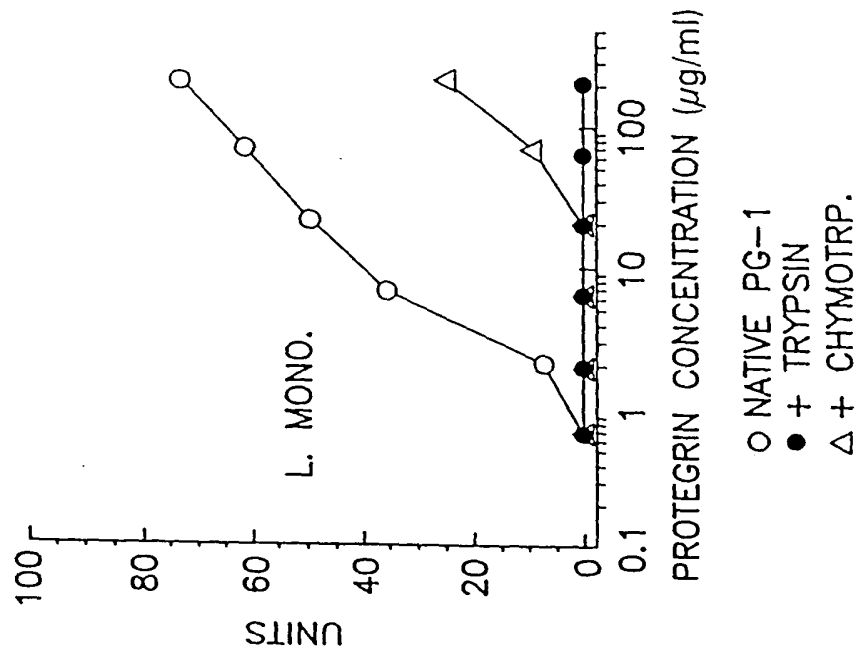


FIG. 12c

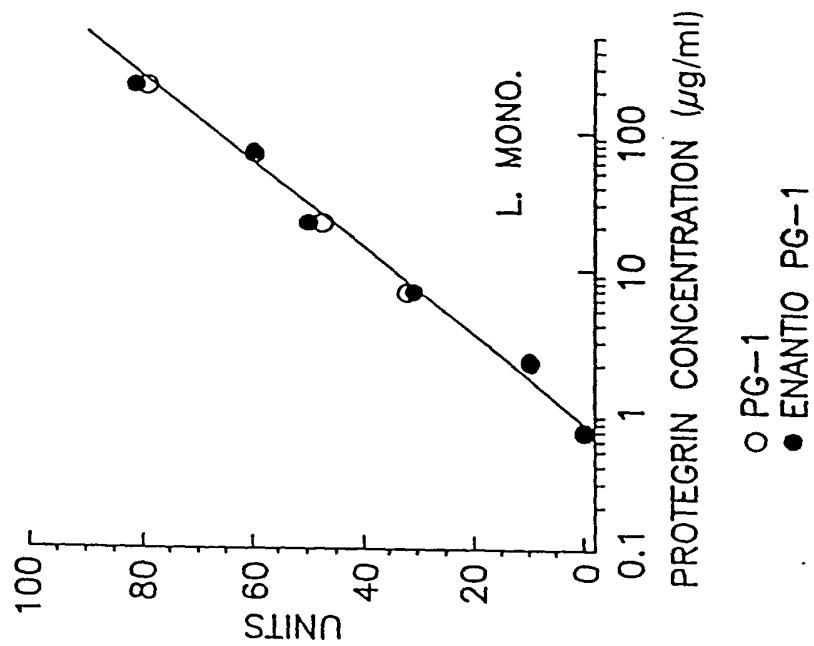


FIG. 12d

OPEN SYMBOLS = KITE, CLOSED SYMBOLS = BULLET

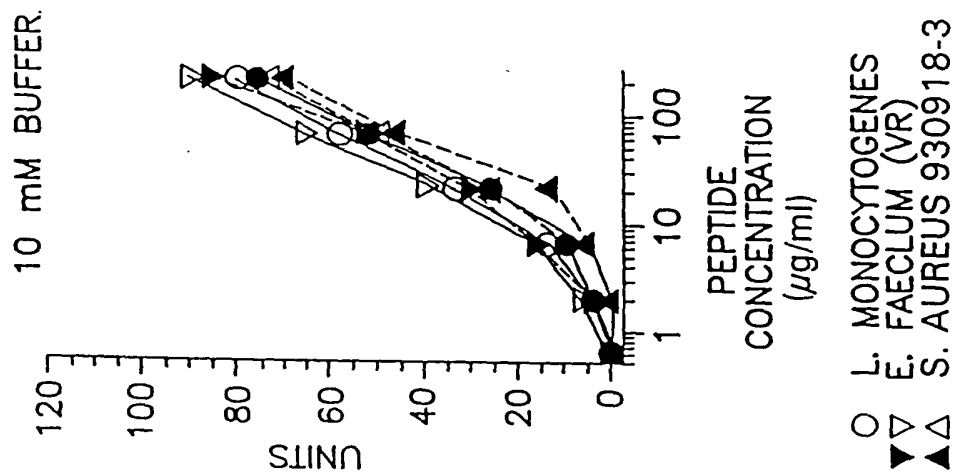


FIG. 13a

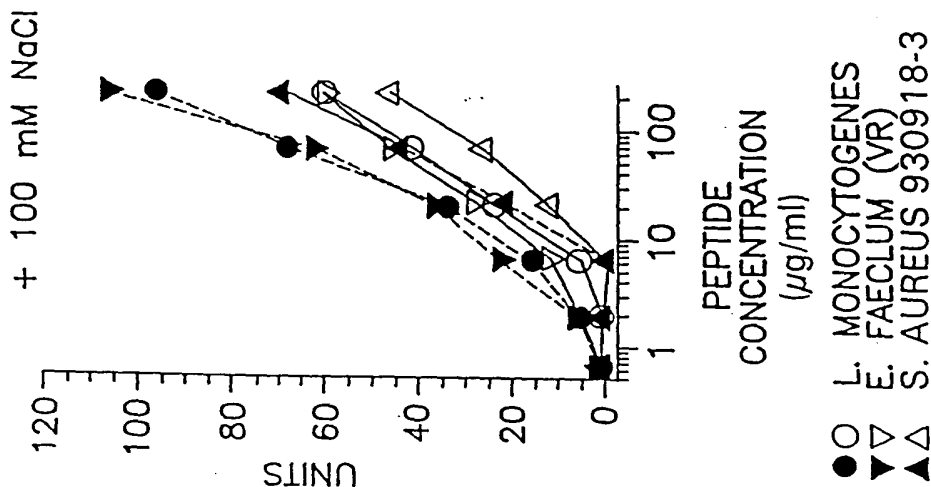


FIG. 13b

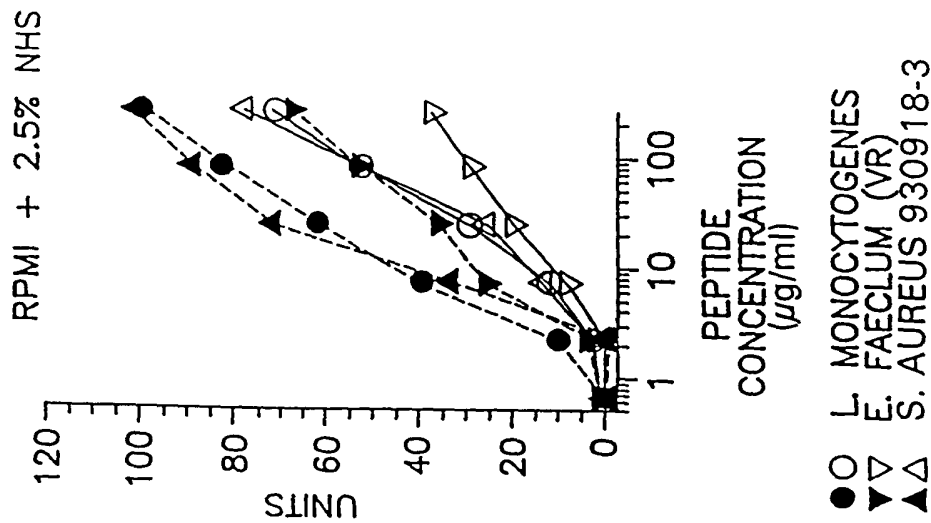


FIG. 13c

OPEN SYMBOLS = KITE, CLOSED SYMBOLS = BULLET

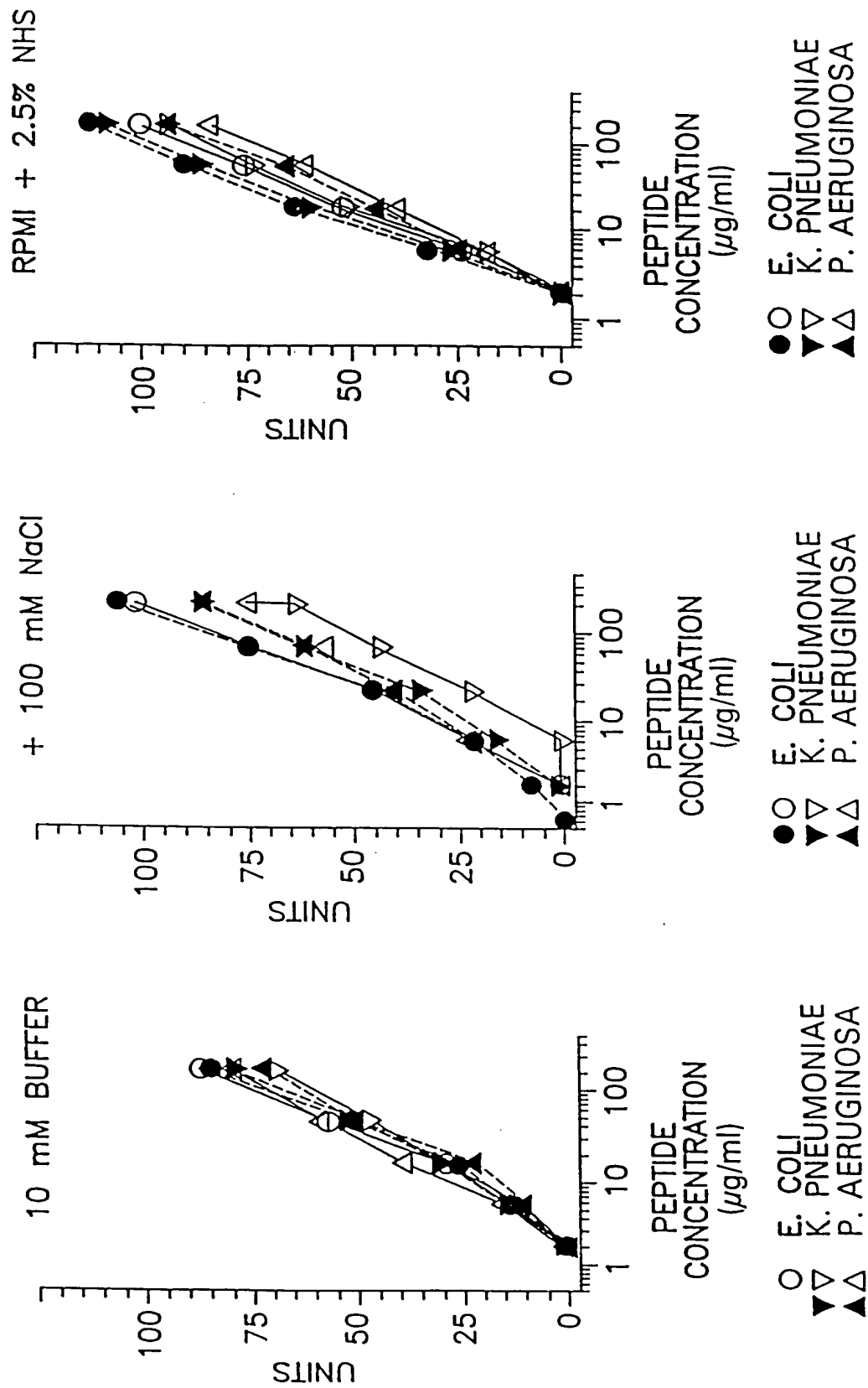


FIG. 14a

FIG. 14b

FIG. 14c

OPEN SYMBOLS = LINEARIZED, CLOSED SYMBOLS = NATIVE

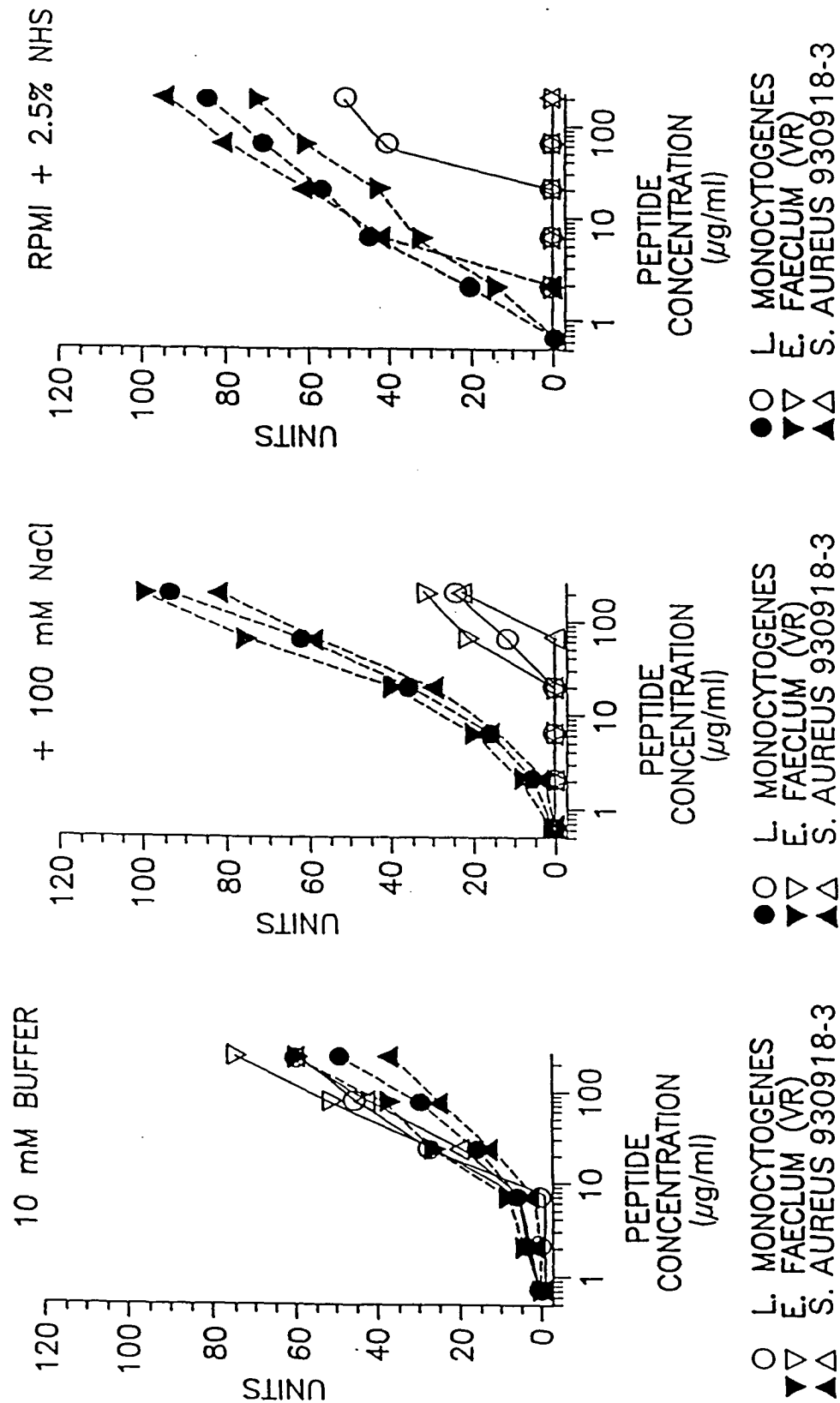


FIG. 15a

FIG. 15b

FIG. 15c

OPEN SYMBOLS = LINEARIZED, CLOSED SYMBOLS = sPG-1

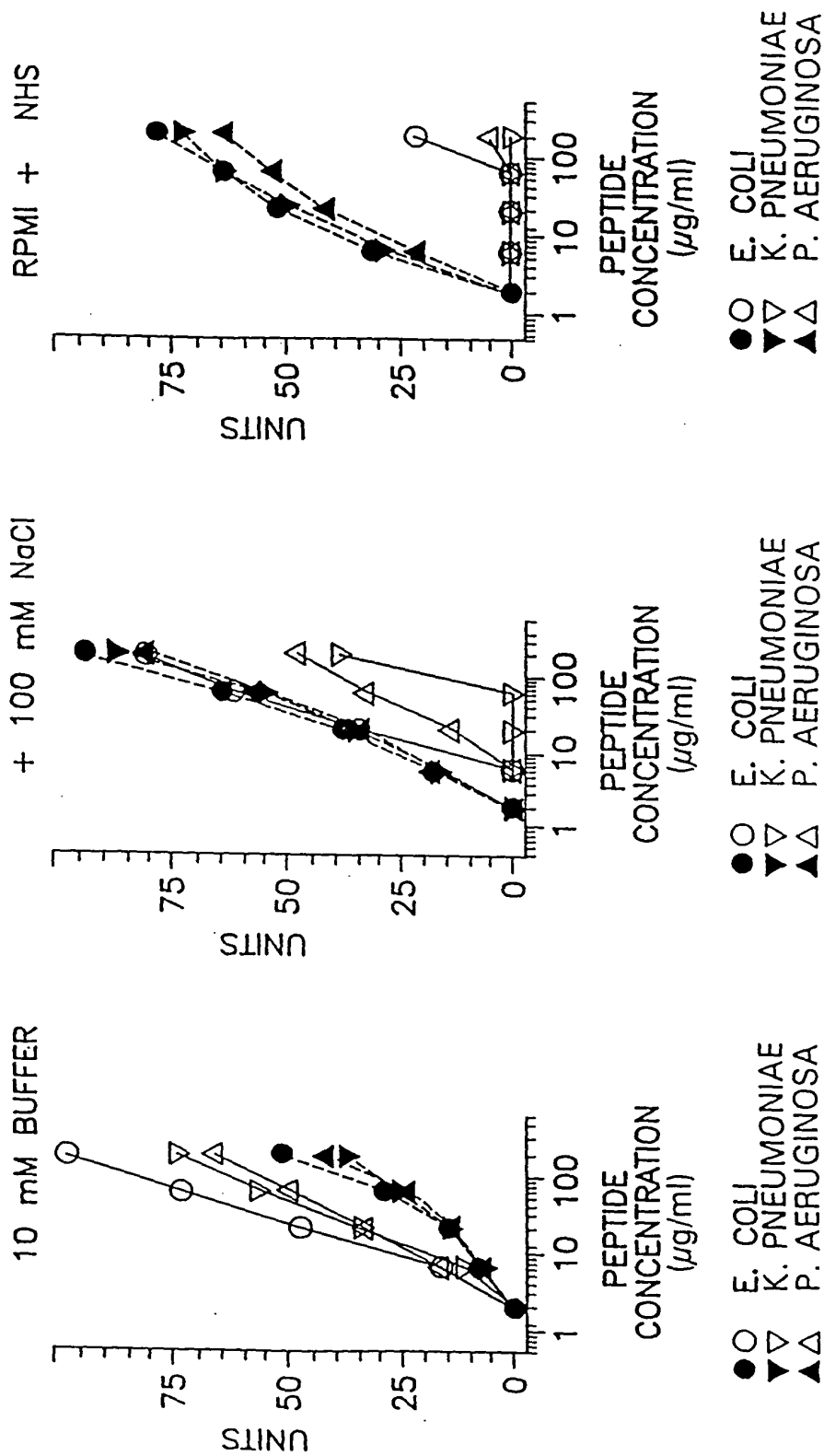


FIG. 16c

FIG. 16b

FIG. 16a